

Figure 1

	Total protein mg	Enzyme activity nkat	Yield %	Specific activity nkat·mg ⁻¹	Purification -fold
Crude extract	1716	51		0.03	
Ammonium sulphate precipitation (30-80%)	890	155	300	0.17	6
DEAE column	10.7	50	98	4.69	158
MonoQ column 60%	1.8	18.6	36	10.23	344
MonoQ column 30%	0.96	6.6	12	6.87	231
Phenylsucrose	0.07	1.2	2.4	18.03	600

nkat = 1nmol caffeoyl-CoA s⁻¹

Figure 2

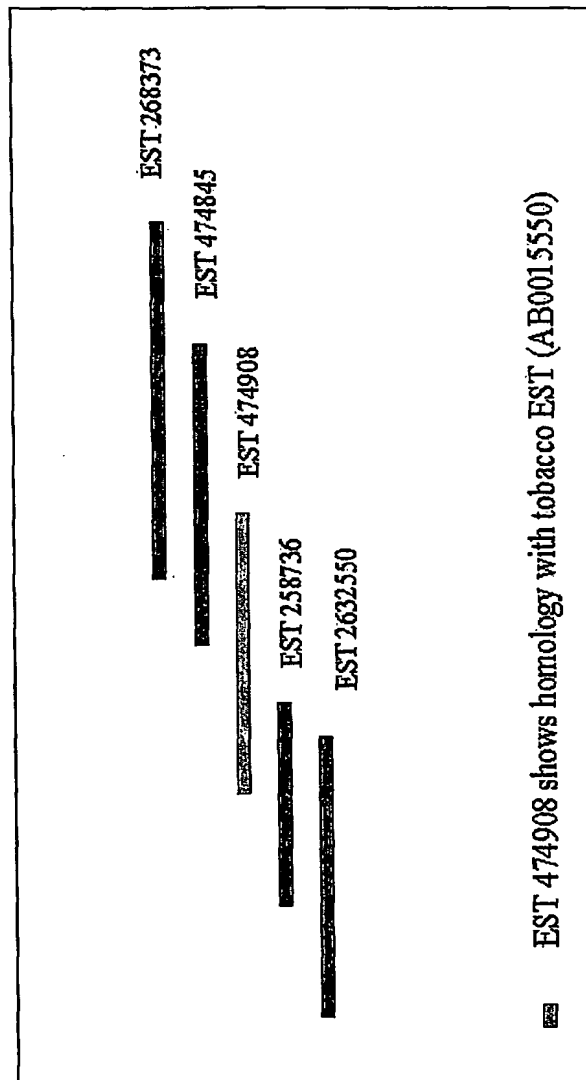


Figure 3

	1					60
tobHQT	MCSEKAMKIN	IKESTLVKPS	KPTPTKELWS	SNLDLIVGRI	HLLTVYFYKE	NCSSNFFLSE
tomHQT	MCSEKAMKIN	IKESTLVKPS	KPTPTKELWS	SNLDLIVGRI	HLLTVYFYKE	NCSSNFFLSE
cons	-----MK--	IK--L-KPS	-PTP--L--	SNLD-I-G--	-----FYK-	-----S-
	61					120
	IMKALSNVI	WSPYPMACRI	APDEQCRIEI	NCNCEGVLEW	EASDSSEWDE	FCDFTP SLEI
	IMKALSNVI	WSPYPMACRI	APDEQCRIEI	NCNCEGVLEW	EASDSSEWDE	FCDFTP SLEI
	--K--LS--L	V--YP-MERL	-R--G--KI	-CN-EG--FV	EA-SD----D	F-D--P---L
	121					180
	RKLLIVVDTG	EDI STFPLII	FOVTRKCCG	WSLGGCVVHT	LSDGLSSITH	INTUSD IARG
	RKLLIVVDTG	EDI STFPLII	FOVTRKCCG	WSLGGCVVHT	LSDGLSSITH	INTUSD IARG
	--L-P-V--S	---T-PL--	-QVT-KCCG	---G--V-H-	--DG-S---F	IN-W--ARG
	181					240
	LSVAIPPFI	RTLLR&RDP	TESFEHWEYH	PPRELISSE	SLESTSPKPS	TTTHIRFESS
	LSVAIPPFI	RTLLR&RDP	TYSEHWEYH	PPRELISSE	SLESTSPKPS	TTTHIRFESS
	-----P--D	R-LL--R-PP	-----H-EY-	P-----	-----	-----K-S--
	241					300
	QLCLRSKSKS	HDCSTYEILN	AHIURCTCKA	RLSDDQLTR	LHWATDGBSE	LCPPLP PGYL
	QLCLRSKSKS	HDCSTYEILN	AHIURCTCKA	RLSDDQLTR	LHWATDGBSE	LCPPLP PGYL
	-L--LK-K--	-----E--	A--VRC--KA	--L-----	L--A--BSR	L-PPLP-GY-
	301					360
	GNVVFICTPM	AKSELLEEF	LTNSKRIHS	ELSHDDNYL	RSALDYLEL	PDLCLIRCH
	GNVVFICTPM	AKSELLEEF	LTNSKRIHS	ELSHDDNYL	RSALDYLEL	PDLCLIRCH
	GN-V-----	---KL---P	L--A-----	A-K-----YL	RS--D--E--	--L-----G-
	361					420
	TYFASNLINI	NSUTBLPVHD	SDFCNGRIH	MCPCILYEC	TYTILPSFNS	KDNLRLAVG
	TYFASNLINI	NSUTBLPVHD	SDFCNGRIH	MCPCILYEC	TYTILPSFNS	KDNLRLAVG
	-----	-SW-R-P--	-DFGWC-P--	-GP-----	-----PS--	-D-----WC
	421	436				
	LDGCHHILFE	KYLYDE				
	LDGCHHILFE	KYLYDE				
	L---M--FR	K-----				

Figure 4

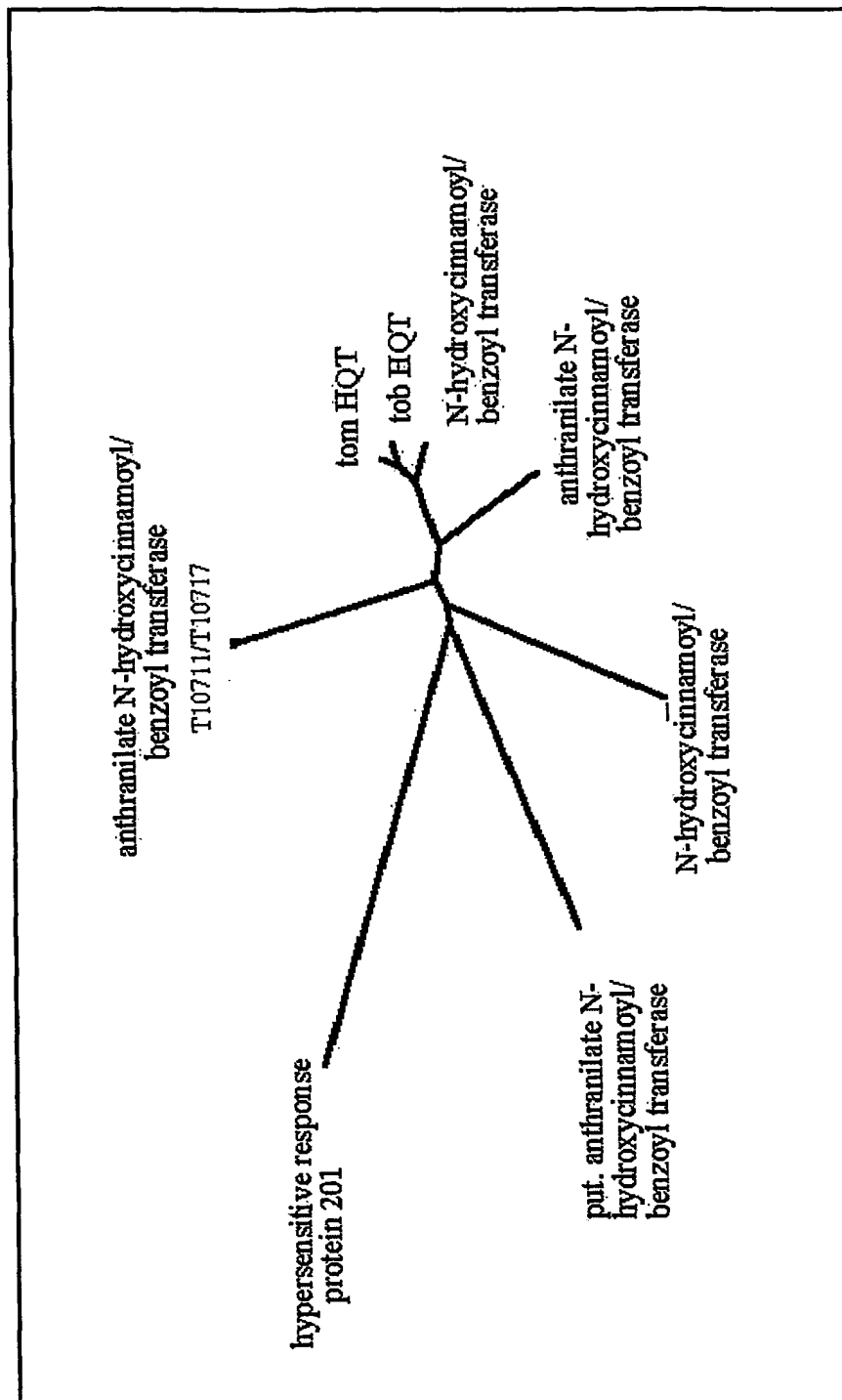


Figure 5

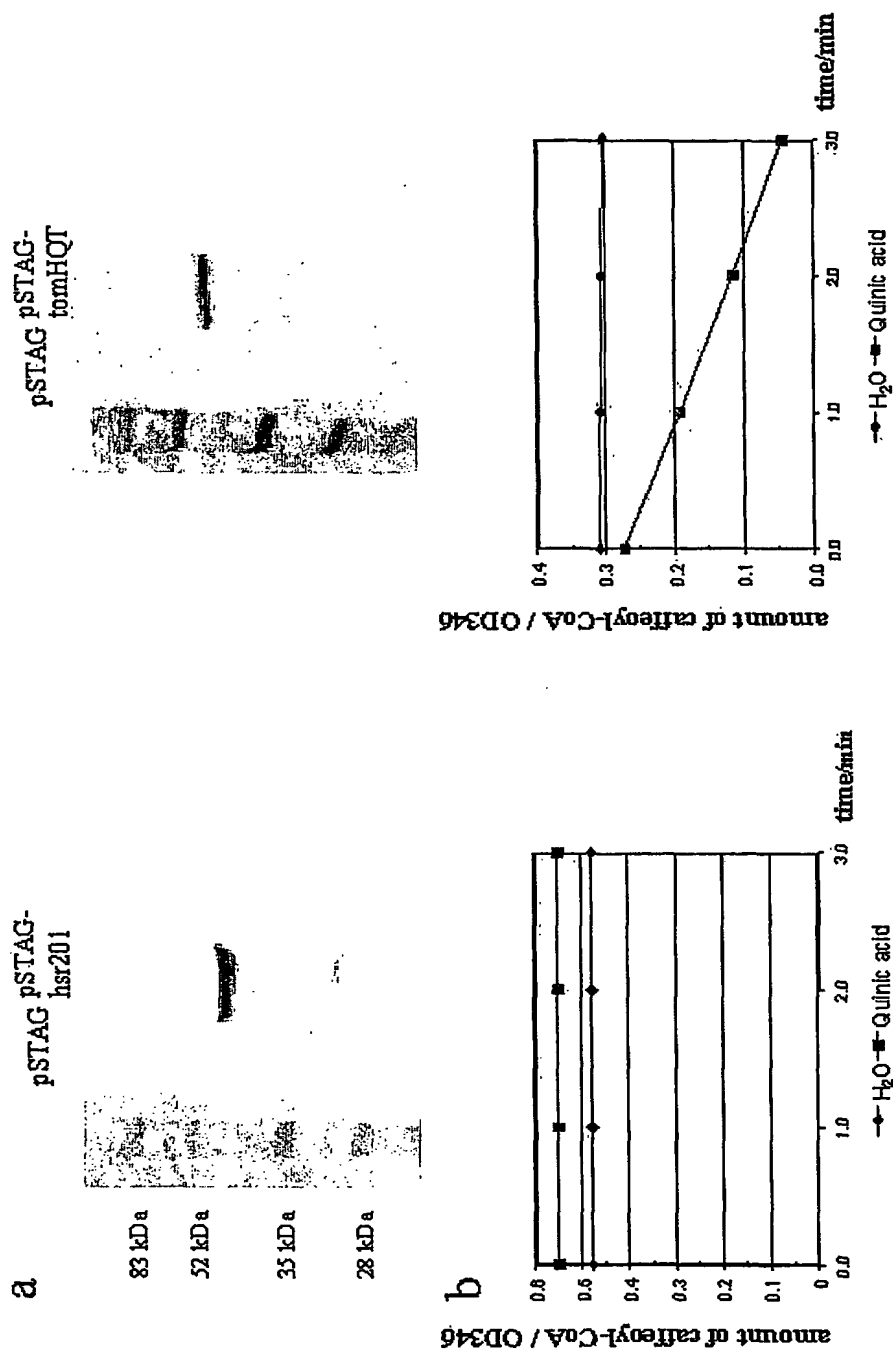


Figure 6

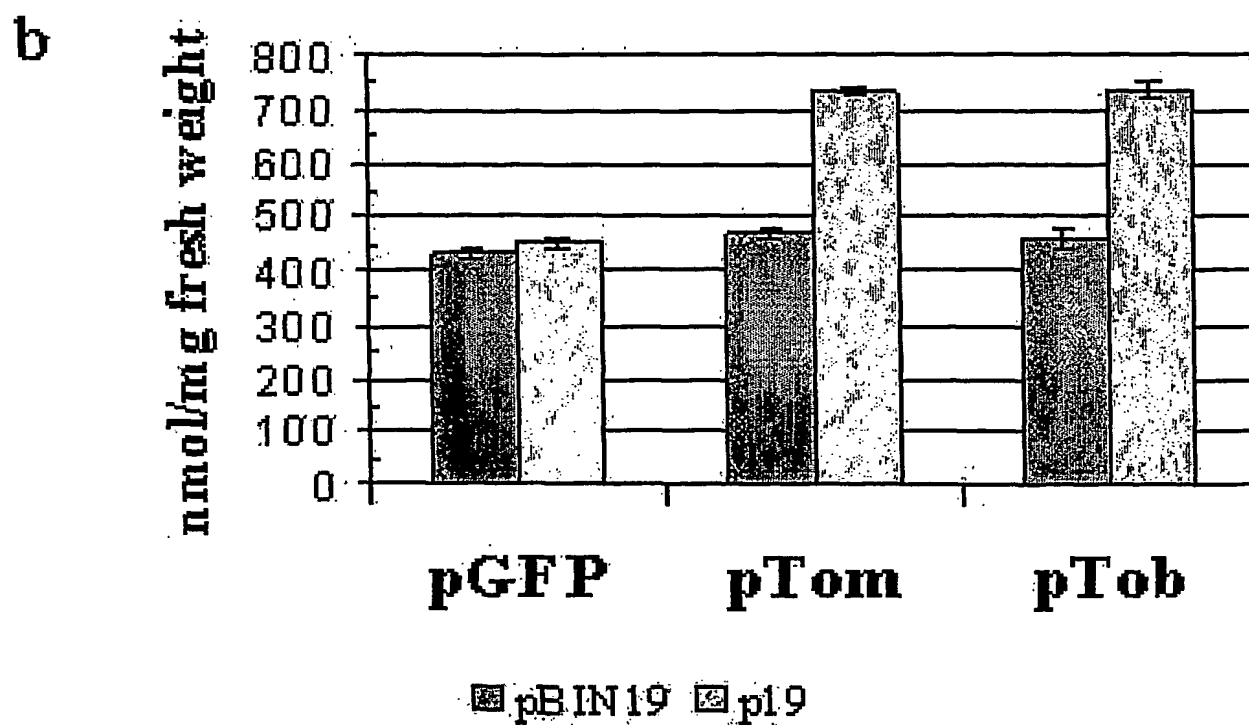
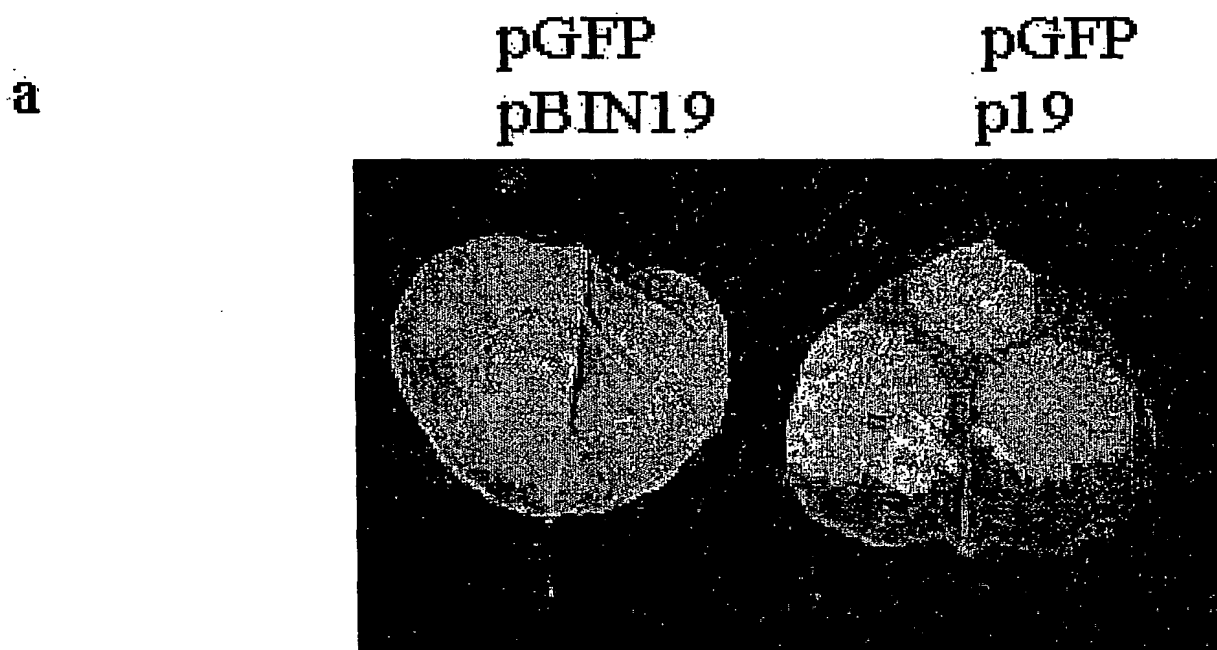


Figure 7a

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TAAACCAAAT GGATCTTCAA ATTTTTTTGA TAATAAAGTT ATTAAAGAAG
CATTAAGTAA TGTTTTAGTT TCATTTTATC CAATGGCTGG AAGATTAGGT
AGGGATGAAC AAGGTAGAAT TGAAGTTAAT TGTAATGGTG AAGGTGTTTT
GTTTGTGAG GCTGAAAGTG ATTCATGTGT TGATGATTTT GGTGATTTTA
CACCATCTTT GGAACCTAGA AAACCTATTC CAAGTGTTGA AACCTCTGGA
GATATCTCAA CTTTCCCACT AGTTATATTT CAGATTACTC GTTCAAGTG
TGGCGGAGTC GCTCTTGGTG GTGGAGTATT CCACACGTTA TCCGATGGTC
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TCCGTGCGAG TCCGCGCGTT CATCGATCGG ACGCTCCTCC GTGCAAGGGA
CCCACCGACA TATTCTTTCG AGCACGTTGA GTACCATCCT cCACCTACCC
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TATACCAAGT CCAAATTCTA AAGATAGGAA CTTGCGTTTG GCTGTTTGTc
TAGATGCTGG TCACATGTCA CTATTTGAAA AATATTTATA TGAATTATGA

Figure 7b

ATGGGAAG TGAAAAATG ATGAAAATTA ATATCAAGGA ATCAACATTA
GTA AACCAT CAAACCAAC ACCAACAAAA AGACTTTGGA GTTCTAACTT
AGATTTAATA GTGGGAAGAA TTCATCTTTT AACAGTATAT TTCTATAAAC
CAAATGGATC TTCAAATTTT TTTGATTCAA AAATAATGAA AGAAGCATT
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CAACATCGTC TTTCGAGCAC GTCGAGTATC ATCCTCCTCC ATCTCTAATT
TCATCATCAA AAAGCTTAGA ATCCACTAGC CCAAAGCCTA GTACCACAAC
CATGTTAAAA TTCTCTAGTG ACCAACTTGG GCTTCTAAAG TCCAAGTCCA
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 AAAAAAAAAA